DT09 Rec'd PCT/PT0 3 0 NOV 2004

## SEQUENCE LISTING

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- <130> D3-A0203P
- <150> JP 2002-161964
- <151> 2002-06-03
- <160> 63
- <170> PatentIn version 3.1
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- <400> 1

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Leu Ala Gly Phe Ala Thr Val Ala Gln Ala Glu Val Lys Leu His Glu

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Leu Ala Leu Leu P	co Leu Leu Phe Thi	Pro Val Thr Lys	Ala Asp Ile
255	260	)	265
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gag ctc acc cag to			
Glu Leu Thr Gln So			Gly Glu Inr
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Val Thr Ile Thr Cy			
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Trp Tyr Gln Arg Ly	's Gln Gly Lys Ser	· Pro Gln Leu Leu	Ile Tyr Gly
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360					300					390					390	
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	gtc															1301
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415 420 425

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1397

Trp Thr Asp Gln Asp Ser Lys Asp Ser Thr Tyr Ser Met Ser Ser Thr

430

435

440

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1445

Leu Thr Leu Thr Lys Asp Glu Tyr Glu Arg His Asn Ser Tyr Thr Cys

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450

455

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35 40 45

Thr Phe Thr Asn Tyr Trp Leu Gly Trp Val Lys Gln Arg Pro Gly His
50 55 60

Gly Leu Glu Trp Ile Gly Asp Ile Tyr Pro Gly Gly Gly Tyr Thr Asn
65 70 75 80

Tyr Asn Glu Lys Phe Lys Gly Lys Ala Thr Leu Thr Ala Asp Thr Ser

85 90 95

Ser Ser Thr Ala Tyr Met Gln Leu Ser Ser Leu Thr Ser Glu Asp Ser

100 105 110

Ala Val Tyr Phe Cys Ala Arg Phe Tyr Tyr Gly Ser Ser Tyr Trp Tyr

115
120 --125

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130 135 140

Thr Thr Pro Pro Ser Val Tyr Pro Leu Ala Pro Gly Ser Ala Ala Gln
145 150 155 160

Thr Asn Ser Met Val Thr Leu Gly Cys Leu Val Lys Gly Tyr Phe Pro
165 170 175

Glu Pro Val Thr Val Thr Trp Asn Ser Gly Ser Leu Ser Ser Gly Val
180 185 190

His Thr Phe Pro Ala Val Leu Gln Ser Asp Leu Tyr Thr Leu Ser Ser 195 200 205

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90

95

85

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115 120 125

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Gln Leu Thr Ser Gly Gly Ala Ser Val Val Cys Phe Leu Asn Asn Phe
145 150 155 160

Tyr Pro Lys Asp Ile Asn Val Lys Trp Lys Ile Asp Gly Ser Glu Arg
165 170 175

Gln Asn Gly Val Leu Asn Ser Trp Thr Asp Gln Asp Ser Lys Asp Ser

180 185 190

Thr Tyr Ser Met Ser Ser Thr Leu Thr Leu Thr Lys Asp Glu Tyr Glu
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70 agggcctggg <210> 14 <211> 70 <212> DNA <213> Artificial <220> <223> a synthetic oligonucleotide for constructing a Fab gene fragment ⟨400⟩ 14 tgcatgagtc agggcctggg ctggtaaggc ctgggacttc agtgaagata tcctgcaagg 60 70 cttctggcta <210> 15 <211> 60 <212> DNA <213> Artificial <220>

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tggtaco	cage agaagecagg acagecacec aaacteetea tetttgetge ateeaaegta 1	80

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35 40 45

Lys Leu Leu Ile Phe Ala Ala Ser Asn Val Glu Ser Gly Val Pro Ala 50 55 60

Arg Phe Ser Gly Ser Gly Ser Gly Thr Asn Phe Ser Leu Asn Ile His

70

75

80

Pro Val Asp Glu Asp Asp Val Ala Met Tyr Phe Cys Gln Gln Ser Arg
85 90 95

Lys Val Pro Tyr Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys Arg

100 110

Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Ser Gln
115 120 125

Val Lys Leu Gln Gln Ser Gly Pro Gly Leu Val Thr Pro Ser Gln Ser 130 135 140

Leu Ser Ile Thr Cys Thr Val Ser Gly Phe Ser Leu Ser Asp Tyr Gly
145 150 155 160

Val His Trp Val Arg Gln Ser Pro Gly Gln Gly Leu Glu Trp Leu Gly
165 170 175

Val Ile Trp Ala Gly Gly Gly Thr Asn Tyr Asn Ser Ala Leu Met Ser 180 185 190

Arg Lys Ser Ile Ser Lys Asp Asn Ser Lys Ser Gln Val Phe Leu Lys
195 200 205

Met Asn Ser Leu Gln Ala Asp Asp Thr Ala Val Tyr Tyr Cys Ala Arg 210 215 220

Asp Lys Gly Tyr Ser Tyr Tyr Tyr Ser Met Asp Tyr Trp Gly Gln Gly
225 230 235 240

Thr Thr Val Thr Val Ser Ser

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cctcag	tcat aatgtccaga ggatctagac cgtagtaaga aaaacttagg gtgaaagttc	120
atcgcg	gccg c	131
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 $\hbox{Met Asp Phe Gln Val Gln Ile Phe Ser Phe Leu Leu Ile Ser Ala Ser}$ 

1 5 10 15 Val Ile Met Ser Arg Gly 20 <210> 47 <211> 70 <212> DNA <213> Artificial <220> <223> an synthetic oligonucleotide for constructing an anti-CD28cst gene fragment <400> 47 tctagagaca tcgagctcac tcagtctcca gcttctttgg ctgtgtctct agggcagaga 60 70 gccaccatct <210> 48 <211> 70 <212> DNA

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acctggcctg		70
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23